

Acta Crystallographica Section E

## Structure Reports

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## 3-Methyl-1,5-diphenyl-4,5-dihydro-1H-pyrazole

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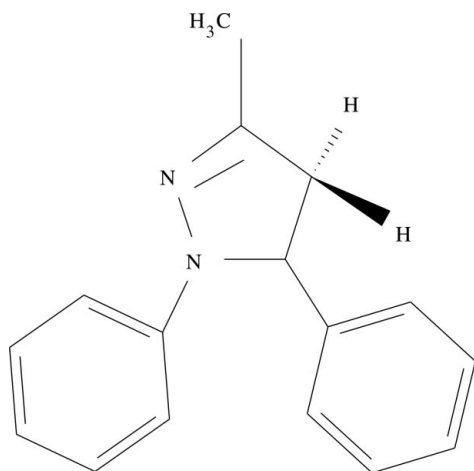
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Key indicators: single-crystal X-ray study;  $T = 301\text{ K}$ ; mean  $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$ ;  $R$  factor = 0.037;  $wR$  factor = 0.131; data-to-parameter ratio = 18.0.

In the title compound,  $\text{C}_{16}\text{H}_{16}\text{N}_2$ , the dihydropyrazole ring adopts a shallow envelope conformation, with the C atom bearing the phenyl group displaced by 0.298 (2)  $\text{\AA}$  from the other atoms (r.m.s. deviation = 0.015  $\text{\AA}$ ). The dihedral angles between the four near coplanar atoms of the central ring and the N- and C-bonded phenyl groups are 13.49 (13) and 82.22 (16)°, respectively.

## Related literature

For background to pyrazoles, see: Govindaraju *et al.* (2012); Jayaroopa *et al.* (2013); Kalirajan *et al.* (2013); Mariappan *et al.* (2010); Shyama *et al.* (2009). For related structures, see: Bakır *et al.* (2011); Fun *et al.* (2011).



## Experimental

## Crystal data

$\text{C}_{16}\text{H}_{16}\text{N}_2$   
 $M_r = 236.31$   
 Monoclinic,  $Cc$   
 $a = 18.1224\text{ (17)\AA}$   
 $b = 7.8055\text{ (6)\AA}$   
 $c = 12.5057\text{ (13)\AA}$   
 $\beta = 132.207\text{ (9)^\circ}$   
 $V = 1310.3\text{ (3)\AA}^3$   
 $Z = 4$   
 Mo  $K\alpha$  radiation  
 $\mu = 0.07\text{ mm}^{-1}$   
 $T = 301\text{ K}$   
 $0.32 \times 0.20 \times 0.18\text{ mm}$

## Data collection

Oxford Diffraction Xcalibur Eos diffractometer  
 11856 measured reflections  
 2973 independent reflections  
 2362 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.033$

## Refinement

$R[F^2 > 2\sigma(F^2)] = 0.037$   
 $wR(F^2) = 0.131$   
 $S = 0.91$   
 2973 reflections  
 165 parameters  
 2 restraints  
 H-atom parameters constrained  
 $\Delta\rho_{\text{max}} = 0.14\text{ e \AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.10\text{ e \AA}^{-3}$

Data collection: *CrysAlis PRO* (Oxford Diffraction, 2009); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *Mercury* (Macrae *et al.*, 2006); software used to prepare material for publication: *Mercury*.

MM thanks the IOE, University of Mysore, for the award of a fellowship. We thank the Solid State and Structural Chemistry Unit, IISc, Bangalore, for the data collection.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB7053).

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## supporting information

*Acta Cryst.* (2013). E69, o602 [doi:10.1107/S1600536813007575]

**3-Methyl-1,5-diphenyl-4,5-dihydro-1H-pyrazole**

**M. Manjula, P. Jayaroopa, B. C. Manjunath, K. Ajay kumar and N. K. Lokanath**

**S1. Comment**

Pyrazoles are five-member heterocycles with two nitrogen atoms in a ring at 1,2-positions. They have been efficiently transformed in to a potential medicinally and pharmaceutically important molecule. Pyrazole derivatives have known to exhibit diverse biological applications such as antidiabetic, anaesthetic, antifungal (Jayaroopa *et al.*, 2013), antiandrogenic, antioxidant, analgesic and anti-inflammatory activities. In addition, they have also showed potential anti-bacterial (Govindaraju *et al.*, 2012), anticancer (Kalirajan *et al.*, 2013), antiamebic, potent and selective inhibitors of tissue-nonspecific alkaline phosphatase (Shyama *et al.*, 2009), anti-inflammatory and protein kinase C inhibitor (Mariappan *et al.*, 2010) properties.

we have synthesized the title compound to study its crystal structure in order to understand the structure-activity relationship for its biological activity.

The title compound C<sub>16</sub> H<sub>16</sub> N<sub>2</sub>, contains two benzene rings (C1-C6 and C8-C13), these two rings are attached to the central pyrazole ring and the molecules are connected by non classical hydrogen bonds. The dihydropyrazole ring is a shallow envelope, with atom C7 displaced from the other four atoms by 0.298 (2) Å. The dihedral angles between the four near coplanar atoms of the central ring and the N- and C-bonded phenyl groups are 13.49 (13) and 82.22 (16)°, respectively. Bond lengths and bond angles are within normal ranges and are comparable to related structures (Baktir *et al.*, 2011 & Fun *et al.*, 2011).

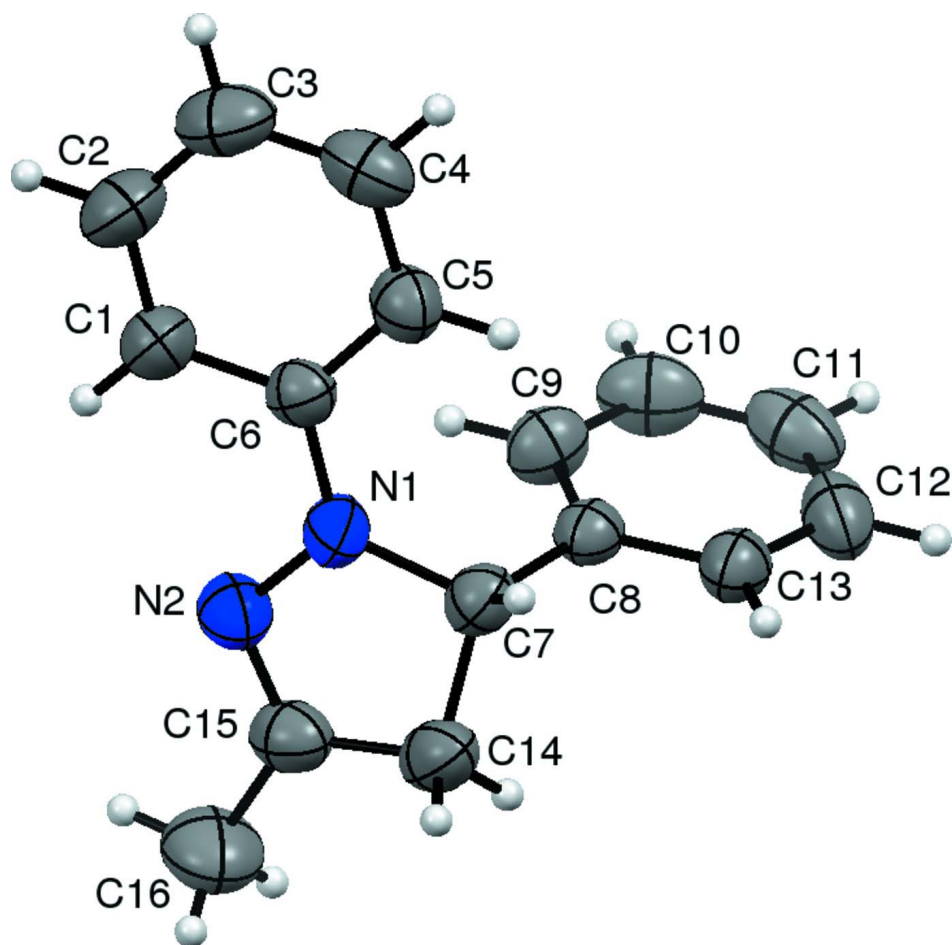
**S2. Experimental**

A mixture of 4-phenylbut-3-en-2-one (0.05 mmol), phenyl hydrazine hydrochloride (0.05 mmol) and sodium acetate (0.05 mmol) in ethyl alcohol (25 ml) was stirred at room temperature for 1 h. The progress of the reaction was monitored by TLC. After the completion of the reaction, the mixture was poured into ice cold water. The solid formed was separated and crystallized with acetonitrile to get the title compound as yellow blocks.

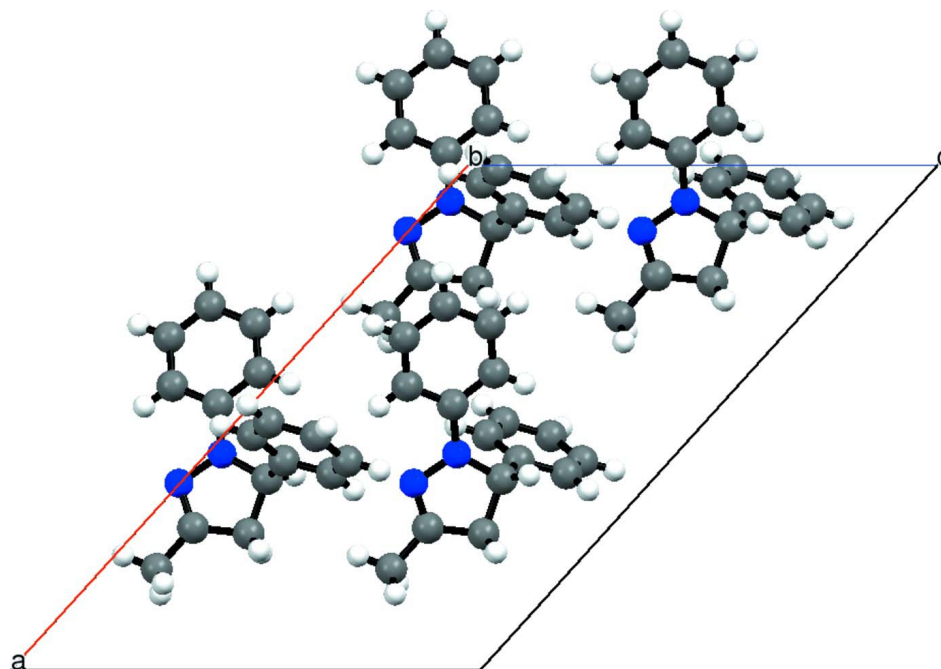
Flash Point: 178 ° C. <sup>1</sup>H NMR (CDCl<sub>3</sub>): δ 2.015 (s, 3H, CH<sub>3</sub>), 2.619–2.671 (q, 1H, C4—H), 3.475–3.536 (q, 1H, C4—H), 5.108–5.146 (q, 1H, C5—H), 6.622–6.655 (t, 1H, Ar—H), 6.805–6.829 (d, 2H, Ar—H), 7.061–7.097 (t, 2H, Ar—H), 7.230–7.373 (m, 5H, Ar—H).

**S3. Refinement**

All hydrogen atoms were located geometrically with C—H = 0.93–0.97) Å and allowed to ride on their parent atoms with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{aromatic C})$ .

**Figure 1**

ORTEP diagram of the title molecule with 50% probability ellipsoids.

**Figure 2**

Packing diagram of molecule, viewed along the crystallographic *b* axis.

### 3-Methyl-1,5-diphenyl-4,5-dihydro-1*H*-pyrazole

#### Crystal data

$C_{16}H_{16}N_2$   
 $M_r = 236.31$   
 Monoclinic, *Cc*  
 Hall symbol: *C* -2yc  
 $a = 18.1224$  (17) Å  
 $b = 7.8055$  (6) Å  
 $c = 12.5057$  (13) Å  
 $\beta = 132.207$  (9)°  
 $V = 1310.3$  (3) Å<sup>3</sup>  
 $Z = 4$

$F(000) = 504$   
 $D_x = 1.198$  Mg m<sup>-3</sup>  
 Melting point = 363–365 K  
 Mo *K*α radiation,  $\lambda = 0.71073$  Å  
 Cell parameters from 2973 reflections  
 $\theta = 3.0$ – $27.6$ °  
 $\mu = 0.07$  mm<sup>-1</sup>  
 $T = 301$  K  
 Block, pale yellow  
 $0.32 \times 0.20 \times 0.18$  mm

#### Data collection

Oxford Diffraction Xcalibur Eos  
 diffractometer  
 Radiation source: fine-focus sealed tube  
 Graphite monochromator  
 Detector resolution: 16.0839 pixels mm<sup>-1</sup>  
 $\omega$  scans  
 11856 measured reflections

2973 independent reflections  
 2362 reflections with  $I > 2\sigma(I)$   
 $R_{int} = 0.033$   
 $\theta_{max} = 27.6$ °,  $\theta_{min} = 3.0$ °  
 $h = -23 \rightarrow 23$   
 $k = -10 \rightarrow 10$   
 $l = -16 \rightarrow 16$

#### Refinement

Refinement on  $F^2$   
 Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.037$   
 $wR(F^2) = 0.131$   
 $S = 0.91$

2973 reflections  
 165 parameters  
 2 restraints  
 Primary atom site location: structure-invariant  
 direct methods

Secondary atom site location: difference Fourier

map

Hydrogen site location: inferred from

neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.1P)^2]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

$$(\Delta/\sigma)_{\max} = 0.001$$

$$\Delta\rho_{\max} = 0.14 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.10 \text{ e } \text{\AA}^{-3}$$

*Special details*

**Geometry.** Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

**Refinement.** Refinement on  $F^2$  for ALL reflections except those flagged by the user for potential systematic errors. Weighted  $R$ -factors  $wR$  and all goodnesses of fit  $S$  are based on  $F^2$ , conventional  $R$ -factors  $R$  are based on  $F$ , with  $F$  set to zero for negative  $F^2$ . The observed criterion of  $F^2 > \sigma(F^2)$  is used only for calculating  $-R$ -factor-obs *etc.* and is not relevant to the choice of reflections for refinement.  $R$ -factors based on  $F^2$  are statistically about twice as large as those based on  $F$ , and  $R$ -factors based on ALL data will be even larger.

*Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
N1	0.57189 (12)	0.5350 (3)	0.53221 (17)	0.0583 (6)
N2	0.63012 (12)	0.5538 (2)	0.49747 (18)	0.0528 (5)
C1	0.43700 (15)	0.4218 (3)	0.2954 (2)	0.0539 (6)
C2	0.34060 (17)	0.3632 (3)	0.1948 (2)	0.0683 (7)
C3	0.27852 (16)	0.3631 (3)	0.2214 (3)	0.0694 (7)
C4	0.31410 (16)	0.4245 (3)	0.3516 (3)	0.0617 (7)
C5	0.41076 (14)	0.4857 (2)	0.4546 (2)	0.0529 (6)
C6	0.47391 (13)	0.4829 (2)	0.42818 (19)	0.0439 (5)
C7	0.62148 (15)	0.5986 (2)	0.6767 (2)	0.0487 (5)
C8	0.58591 (12)	0.7740 (2)	0.67789 (18)	0.0425 (5)
C9	0.53469 (16)	0.8844 (3)	0.5615 (2)	0.0558 (6)
C10	0.50989 (18)	1.0475 (3)	0.5715 (3)	0.0710 (8)
C11	0.53636 (18)	1.1018 (3)	0.6979 (3)	0.0725 (9)
C12	0.58650 (17)	0.9929 (3)	0.8138 (3)	0.0644 (8)
C13	0.61057 (14)	0.8294 (3)	0.80360 (19)	0.0502 (6)
C14	0.73007 (16)	0.6028 (3)	0.7433 (2)	0.0618 (6)
C15	0.71843 (15)	0.5947 (2)	0.6134 (2)	0.0563 (6)
C16	0.8000 (2)	0.6260 (4)	0.6152 (4)	0.0878 (11)
H1	0.47760	0.42050	0.27480	0.0650*
H2	0.31700	0.32270	0.10660	0.0820*
H3	0.21360	0.32240	0.15260	0.0830*
H4	0.27280	0.42500	0.37110	0.0740*
H5	0.43330	0.52860	0.54160	0.0640*
H7	0.61430	0.51580	0.72810	0.0580*
H9	0.51670	0.84890	0.47570	0.0670*
H10	0.47520	1.12070	0.49240	0.0850*
H11	0.52030	1.21200	0.70480	0.0870*
H12	0.60430	1.02910	0.89940	0.0770*
H13	0.64380	0.75570	0.88220	0.0600*
H14A	0.76320	0.70760	0.79730	0.0740*

H14B	0.76730	0.50530	0.80630	0.0740*
H16A	0.77370	0.62300	0.51830	0.1320*
H16B	0.85000	0.53900	0.67170	0.1320*
H16C	0.82900	0.73640	0.65690	0.1320*

*Atomic displacement parameters ( $\text{\AA}^2$ )*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
N1	0.0465 (9)	0.0796 (11)	0.0514 (9)	−0.0147 (8)	0.0339 (8)	−0.0253 (8)
N2	0.0518 (9)	0.0527 (8)	0.0596 (9)	−0.0064 (7)	0.0397 (8)	−0.0083 (7)
C1	0.0477 (11)	0.0618 (11)	0.0449 (9)	0.0008 (8)	0.0281 (9)	−0.0037 (9)
C2	0.0521 (12)	0.0800 (14)	0.0474 (11)	−0.0079 (10)	0.0231 (10)	−0.0109 (10)
C3	0.0457 (11)	0.0665 (13)	0.0668 (14)	−0.0070 (9)	0.0259 (11)	−0.0034 (10)
C4	0.0532 (11)	0.0519 (10)	0.0866 (15)	0.0036 (9)	0.0497 (11)	0.0054 (10)
C5	0.0551 (11)	0.0494 (10)	0.0590 (11)	−0.0021 (8)	0.0403 (10)	−0.0077 (8)
C6	0.0413 (9)	0.0383 (7)	0.0461 (9)	0.0022 (7)	0.0269 (8)	−0.0015 (7)
C7	0.0479 (9)	0.0476 (9)	0.0425 (9)	−0.0016 (7)	0.0271 (8)	−0.0040 (7)
C8	0.0391 (8)	0.0454 (8)	0.0420 (8)	−0.0051 (7)	0.0269 (7)	−0.0026 (7)
C9	0.0574 (11)	0.0567 (11)	0.0491 (10)	0.0004 (8)	0.0341 (9)	0.0071 (8)
C10	0.0650 (14)	0.0544 (11)	0.0800 (16)	0.0095 (10)	0.0431 (13)	0.0199 (11)
C11	0.0664 (14)	0.0499 (12)	0.110 (2)	0.0015 (9)	0.0628 (15)	−0.0042 (12)
C12	0.0686 (14)	0.0652 (13)	0.0789 (15)	−0.0099 (10)	0.0575 (13)	−0.0201 (11)
C13	0.0490 (10)	0.0567 (10)	0.0457 (10)	−0.0039 (8)	0.0322 (9)	−0.0032 (8)
C14	0.0480 (11)	0.0595 (11)	0.0545 (11)	0.0037 (9)	0.0249 (9)	−0.0119 (9)
C15	0.0463 (11)	0.0505 (10)	0.0644 (12)	−0.0024 (8)	0.0340 (10)	−0.0060 (9)
C16	0.0631 (15)	0.105 (2)	0.100 (2)	−0.0193 (14)	0.0567 (15)	−0.0180 (16)

*Geometric parameters ( $\text{\AA}$ ,  $^\circ$ )*

N1—N2	1.393 (4)	C14—C15	1.492 (3)
N1—C6	1.381 (3)	C15—C16	1.484 (6)
N1—C7	1.462 (3)	C1—H1	0.9300
N2—C15	1.284 (3)	C2—H2	0.9300
C1—C2	1.375 (4)	C3—H3	0.9300
C1—C6	1.393 (3)	C4—H4	0.9300
C2—C3	1.371 (5)	C5—H5	0.9300
C3—C4	1.373 (4)	C7—H7	0.9800
C4—C5	1.387 (4)	C9—H9	0.9300
C5—C6	1.388 (4)	C10—H10	0.9300
C7—C8	1.518 (3)	C11—H11	0.9300
C7—C14	1.538 (4)	C12—H12	0.9300
C8—C9	1.381 (3)	C13—H13	0.9300
C8—C13	1.384 (3)	C14—H14A	0.9700
C9—C10	1.383 (4)	C14—H14B	0.9700
C10—C11	1.375 (4)	C16—H16A	0.9600
C11—C12	1.371 (4)	C16—H16B	0.9600
C12—C13	1.382 (4)	C16—H16C	0.9600

N2—N1—C6	120.09 (17)	C2—C3—H3	121.00
N2—N1—C7	112.6 (2)	C4—C3—H3	121.00
C6—N1—C7	126.7 (2)	C3—C4—H4	120.00
N1—N2—C15	107.8 (2)	C5—C4—H4	119.00
C2—C1—C6	120.1 (3)	C4—C5—H5	120.00
C1—C2—C3	121.6 (2)	C6—C5—H5	120.00
C2—C3—C4	118.6 (3)	N1—C7—H7	110.00
C3—C4—C5	121.0 (3)	C8—C7—H7	110.00
C4—C5—C6	120.2 (2)	C14—C7—H7	110.00
N1—C6—C1	120.5 (3)	C8—C9—H9	120.00
N1—C6—C5	121.00 (18)	C10—C9—H9	120.00
C1—C6—C5	118.5 (2)	C9—C10—H10	120.00
N1—C7—C8	113.95 (16)	C11—C10—H10	120.00
N1—C7—C14	100.2 (2)	C10—C11—H11	120.00
C8—C7—C14	111.89 (18)	C12—C11—H11	120.00
C7—C8—C9	122.80 (19)	C11—C12—H12	120.00
C7—C8—C13	118.56 (16)	C13—C12—H12	120.00
C9—C8—C13	118.55 (19)	C8—C13—H13	120.00
C8—C9—C10	120.5 (2)	C12—C13—H13	120.00
C9—C10—C11	120.3 (2)	C7—C14—H14A	111.00
C10—C11—C12	119.7 (2)	C7—C14—H14B	111.00
C11—C12—C13	120.1 (3)	C15—C14—H14A	111.00
C8—C13—C12	120.8 (2)	C15—C14—H14B	111.00
C7—C14—C15	102.47 (18)	H14A—C14—H14B	109.00
N2—C15—C14	113.4 (3)	C15—C16—H16A	109.00
N2—C15—C16	122.1 (2)	C15—C16—H16B	109.00
C14—C15—C16	124.5 (2)	C15—C16—H16C	109.00
C2—C1—H1	120.00	H16A—C16—H16B	110.00
C6—C1—H1	120.00	H16A—C16—H16C	110.00
C1—C2—H2	119.00	H16B—C16—H16C	109.00
C3—C2—H2	119.00		
C6—N1—N2—C15	−176.91 (18)	C4—C5—C6—N1	−176.63 (19)
C7—N1—N2—C15	11.2 (2)	C4—C5—C6—C1	1.7 (3)
N2—N1—C6—C1	10.6 (3)	N1—C7—C8—C9	−18.0 (4)
N2—N1—C6—C5	−171.14 (17)	N1—C7—C8—C13	165.6 (2)
C7—N1—C6—C1	−178.8 (2)	C14—C7—C8—C9	94.8 (3)
C7—N1—C6—C5	−0.5 (3)	C14—C7—C8—C13	−81.6 (3)
N2—N1—C7—C8	101.5 (3)	N1—C7—C14—C15	17.20 (19)
N2—N1—C7—C14	−18.2 (2)	C8—C7—C14—C15	−103.93 (19)
C6—N1—C7—C8	−69.7 (3)	C7—C8—C9—C10	−175.5 (3)
C6—N1—C7—C14	170.6 (2)	C13—C8—C9—C10	0.8 (4)
N1—N2—C15—C14	1.8 (2)	C7—C8—C13—C12	175.1 (3)
N1—N2—C15—C16	−179.6 (2)	C9—C8—C13—C12	−1.4 (4)
C6—C1—C2—C3	0.1 (3)	C8—C9—C10—C11	0.3 (5)
C2—C1—C6—N1	177.2 (2)	C9—C10—C11—C12	−0.8 (6)
C2—C1—C6—C5	−1.1 (3)	C10—C11—C12—C13	0.2 (6)
C1—C2—C3—C4	0.5 (4)	C11—C12—C13—C8	0.9 (5)

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C2—C3—C4—C5	0.1 (4)	C7—C14—C15—N2	−12.9 (2)
C3—C4—C5—C6	−1.1 (3)	C7—C14—C15—C16	168.6 (2)

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